

# Alignment View Print

	1	10		20		30																			
seq2 seq	M	E	G	D	G	R	L	V	L	L	L														
seq26	M	E	G	D	G	R	L	V	L	L	L														
Consensus	M	E	G	D	G	R	L	V	L	L	L														
	40		50		60																				
seq2 seq	G	A	C	Y	A	A	L	C	K	E	D	E	V	G	E	C	C	K	C	C	C				
seq26	G	A	C	Y	A	A	L	C	K	E	D	E	V	G	E	C	C	K	C	C	C				
Consensus	G	A	C	Y	A	A	L	C	K	E	D	E	V	G	E	C	C	K	C	C	C				
	70		80		90																				
seq2 seq	R	V	K	E	A	C	G	E	L	G	V	C	E	C	C	Y	I	A	H	L	N	G	L		
seq26	R	V	K	E	A	C	G	E	L	G	V	C	E	C	C	Y	I	A	H	L	N	G	L		
Consensus	R	V	K	E	A	C	G	E	L	G	V	C	E	C	C	Y	I	A	H	L	N	G	L		
	100		110		120																				
seq2 seq	K	C	L	Q	C	Q	M	C	D	A	M	G	L	R	A	R	N	C	R	E	N	A	V	C	G
seq26	K	C	L	Q	C	Q	M	C	D	A	M	G	L	R	A	R	N	C	R	E	N	A	V	C	G
Consensus	K	C	L	Q	C	Q	M	C	D	A	M	G	L	R	A	R	N	C	R	E	N	A	V	C	G
	130		140		150																				
seq2 seq	C	G	H	F	C	I	V	Q	D	G	D	H	C	A	A	C	R	A	A	A	A	A	A	A	
seq26	C	G	H	F	C	I	V	Q	D	G	D	H	C	A	A	C	R	A	A	A	A	A	A	A	
Consensus	C	G	H	F	C	I	V	Q	D	G	D	H	C	A	A	C	R	A	A	A	A	A	A	A	
	160		170		180																				
seq2 seq	Q	K	G	G	E	Q	D	L	C	Q	N	C	G	L	E	E	C	Q	N	G	L	E	E	C	Q
seq26	Q	K	G	G	E	Q	D	L	C	Q	N	C	G	L	E	E	C	Q	N	G	L	E	E	C	Q
Consensus	Q	K	G	G	E	Q	D	L	C	Q	N	C	G	L	E	E	C	Q	N	G	L	E	E	C	Q
	190		200		210																				
seq2 seq	H	Q	K	C	W	L	V	K	A	G	A	G	H	W	V	W	W	F	L	G	L	L	L	L	
seq26	H	Q	K	C	W	L	V	K	A	G	A	G	H	W	V	W	W	F	L	G	L	L	L	L	
Consensus	H	Q	K	C	W	L	V	K	A	G	A	G	H	W	V	W	W	F	L	G	L	L	L	L	

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	220	230	240
seq2 seq	V L V L V C V C	L L C V K R R K	R C D V V K V L V
seq26	V L V L V C V C	L L C V K R R K	R C D V V K V L V
Consensus	V L V L V C V C	L L C V K R R K	R C D V V K V L V
	250	260	270
seq2 seq	V C R K R C E A E C	E A V I E A L O A	D V V A V E
seq26	V C R K R C E A E C	E A V I E A L O A	D V V A V E
Consensus	V C R K R C E A E C	E A V I E A L O A	D V V A V E
	280	290	300
seq2 seq	E L L L L G R	N H	
seq26	E L L L L G R	N H	
Consensus	E L L L L G R	N H	

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